

RAW SEQUENCE LISTING

**ERROR REPORT** 

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

JUL 2 4 2006

Application Serial Number:

Source:

Date Processed by STIC:

10/508,759

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE-LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
- Through 2.4 U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
  - Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
     U.S. Patent and Trademark Office, 220 20<sup>th</sup> Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

## Raw Sequence Listing Error Summary

RROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/5/08/15 PRADEMARKS
TTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
IWrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5 <sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:  (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing.  Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
0Invalid <213> Response - "	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.  Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
3 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

AMC - Biotechnology Systems Branch - 09/09/2003





PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/508,759

DATE: 09/29/2004

-------

Input Set : A:\30215APG SEQ.txt

TIME: 16:59:56

Output Set: N:\CRF4\09292004\J508759.raw

```
3 <110> APPLICANT: APROGEN INC.
      5 <120> TITLE OF INVENTION: HUMANIZED ANTIBODY AND PROCESS FOR PREPARING SAME
     7 <130> FILE REFERENCE: PCA30215/APG
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/508,759
C--> 9 <141> CURRENT FILING DATE: 2004-09-22
     9 <150> PRIOR APPLICATION NUMBER: KR10-2002-0015708
     10 <151> PRIOR FILING DATE: 2002-03-22
     12 <160> NUMBER OF SEQ ID NOS: 38
     14 <170> SOFTWARE: KopatentIn 1.71
     16 <210> SEQ ID NO: 1
                                                                     Does Not Comply
     17 <211> LENGTH: 345
                                                                Corrected Diskette Needec
     18 <212> TYPE: DNA
     19 <213 > ORGANISM: Artificial Sequence
     21 <220> FEATURE:
     22 <223> OTHER INFORMATION: HEAVY CHAIN of
     24 <400> SEQUENCE: 1
     25 caggtccagc tggtgcagtc tggagctgaa gtgaagaagc ctggggcctc agtgaaggtt
                                                                                  120
    27 tectgeaaag ettetggeta cacetteace agtgettgga tgaactgggt gegacaggee
     29 cctggacagg gtcttgagtg gatgggacgg atttatccta gtggtggaag cactagctac
                                                                                  180
                                                                                  240
     31 gcacagaagt tccagggcag agtcacaatg actgcagaca aatccacgag cacagtctac
     33 atggagetea geageetgag atetgaggae aeggeggtgt attactgtge aagagagtae
                                                                                  300
     35 cgggttgccc gttggggcca aggaactctg gtcactgtct cttca
     38 <210> SEQ ID NO: 2
     39 <211> LENGTH: 115
     40 <212> TYPE: PRT
     41 <213 > ORGANISM: Artificial Sequence
     43 <220> FEATURE:
     44.<223> OTHER INFORMATION: HEAVY CHAIN of HZVII
     47 <400> SEQUENCE: 2
     48 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Ala Pro Gly Ala
```

48 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Ala Pro Gly Ala
49 1 5 10 15

51 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Ala

52 20 25 30 54 Trp Met Asn Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met

54 TIP MEC ASIT TIP VAL AIG GIT AIR TIO GIF GIT GIF DEC GIR IIP NO.

57 Gly Arg Ile Tyr Pro Ser Gly Gly Ser Thr Ser Tyr Ala Gln Lys Phe

58 50 55 60

60 Gln Gly Arg Val Thr Met Thr Ala Asp Lys Ser Thr Ser Thr Val Tyr 61 65 70 75 80

61 65 70 75 80 63 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys

63 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp inn Ala val lyn lyn cys 64 85 90 95

66 Ala Arg Glu Tyr Arg Val Ala Arg Trp Gly Gln Gly Thr Leu Val Thr 67 100 105 110 RAW SEQUENCE LISTING DATE: 09/29/2004
PATENT APPLICATION: US/10/508,759 TIME: 16:59:56

Input Set : A:\30215APG SEQ.txt

```
69 Val Ser Ala
70
           115
73 <210> SEQ ID NO: 3
74 <211> LENGTH: 336
75 <212> TYPE: DNA
76 <213> ORGANISM: Artificial Sequence
78 <220> FEATURE:
79 <223> OTHER INFORMATION: LIGHT CHAIN of (HZVII
-82 <400> SEQUENCE: 3
83 gatategtga tgacceaaac tecaetttet ttgteggtta eeeetggaca accageetet
                                                                               60
85 atctcttgca agtcaagtca gagcctctta tatagtaatg gaaaaaccta tttgaattgg
                                                                              120
87 ttattacaga agccaggcca gcctccacag cgcctaatct atctggtgtc taatcgggac
                                                                              180
                                                                              240
89 tctqqaqtcc ctqacaggtt cagtggcagt ggatcaggaa cagattttac actgaaaatc
91 agcagagtgg aggctgagga tgttggagtt tattactgcg tgcaaggtac acattttcct
                                                                              300
                                                                              336
93 cagacgttcg gtggaggcac caaggtggaa atcaaa
96 <210> SEQ ID NO: 4
97 <211> LENGTH: 112
98 <212> TYPE: PRT
99 <213> ORGANISM: Artificial Sequence
101 <220> FEATURE:
102 <223 > OTHER INFORMATION: LIGHT CHAIN of (HZVII
105 <400> SEQUENCE: 4
106 Asp Ile Val Met Thr Gln Thr Pro Leu Ser Leu Ser Val Thr Pro Gly
                                          10
                       5
109 Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu Leu Tyr Ser
                                                          30
                  20
110
112 Asn Gly Lys Thr Tyr Leu Asn Trp Leu Leu Gln Lys Pro Gly Gln Pro
113
             35
115 Pro Gln Arg Leu Ile Tyr Leu Val Ser Asn Arg Asp Ser Gly Val Pro
                              55
116
         50
118 Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
                          70
121 Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys Val Gln Gly
                      85
124 Thr His Phe Pro Gln Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
                                     105
125
                 100
130 <210> SEQ ID NO: 5
131 <211> LENGTH: 26
132 <212> TYPE: DNA
133 <213> ORGANISM: Artificial Sequence
135 <220> FEATURE:
136 <223> OTHER INFORMATION Ryu94
139 <400> SEQUENCE: 5
                                                                                26
140 qaqaattcac attcacgatg tacttg
143 <210> SEQ ID NO: 6
144 <211> LENGTH: 33
145 <212> TYPE: DNA
146 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
```

RAW SEQUENCE LISTING DATE: 09/29/2004
PATENT APPLICATION: US/10/508,759 TIME: 16:59:56

Input Set : A:\30215APG\_SEQ.txt

```
149 <223> OTHER INFORMATION:
                              HUR43-1
152 <400> SEQUENCE: 6
153 ctgctgcagc tggacctgac tctggacacc att
                                                                                 33
156 <210> SEQ ID NO: 7
157 <211> LENGTH: 33
158 <212> TYPE: DNA
159 <213> ORGANISM: Artificial Sequence
161 <220> FEATURE:
162 <223> OTHER INFORMATION(:
                              HUR44-1
165 <400> SEQUENCE: 7
                                                                                 33
166 caggtccagc tgcagcagtc tggacctgaa ctg
.169 <210> SEQ ID NO: 8
170 <211> LENGTH: 33
171 <212> TYPE: DNA
172 <213> ORGANISM: Artificial Sequence
174 <220> FEATURE:
175 <223> OTHER INFORMATION
                              HUR45
178 <400> SEQUENCE: 8
                                                                                 33
179 tgggcccttg gtggaggctg cagagacagt gac
182 <210> SEQ ID NO: 9
183 <211> LENGTH: 33
184 <212> TYPE: DNA
185 <213> ORGANISM: Artificial Sequence
187 <220> FEATURE:
                              HUR46-1
188 <223> OTHER INFORMATION
191 <400> SEOUENCE: 9
                                                                                 33
192 gcctccacca agggcccatc ggtcttcccc ctg
195 <210> SEQ ID NO: 10
196 <211> LENGTH: 28
197 <212> TYPE: DNA
198 <213> ORGANISM: Artificial Sequence
200 <220> FEATURE:
201 <223> OTHER INFORMATION
                              HUR31
204 <400> SEQUENCE: 10
                                                                                 28
205 cageggeege teatttacce ggggacag
208 <210> SEQ ID NO: 11
209 <211> LENGTH: 26
210 <212> TYPE: DNA
211 <213 > ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223 > OTHER INFORMATION
217 <400> SEQUENCE: 11
                                                                                 26
218 caaagcttgg aagcaagatg gattca
221 <210> SEQ ID NO: 12
222 <211> LENGTH: 27
223 <212> TYPE: DNA
224 <213> ORGANISM: Artificial Sequence
226 <220> FEATURE:
227 <223> OTHER INFORMATION:
                              HUR48
```

RAW SEQUENCE LISTING DATE: 09/29/2004
PATENT APPLICATION: US/10/508,759 TIME: 16:59:56

Input Set : A:\30215APG SEQ.txt

```
230 <400> SEOUENCE: 12
231 caagatatcc ccacaggtac cagatac
                                                                                 27
234 <210> SEQ ID NO: 13
235 <211> LENGTH: 27
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION
                              HUR49
.243 <400> SEQUENCE: 13
                                                                                 27
244 tgtggggata tcttgatgac ccaaact
247 <210> SEQ ID NO: 14
248 <211> LENGTH: 27
249 <212> TYPE: DNA
250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION ( HUR50
256 <400> SEQUENCE: 14
                                                                                27
257 cacagatett ttgattteca gettggt
260 <210> SEQ ID NO: 15
261 <211> LENGTH: 27
262 <212> TYPE: DNA
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <223> OTHER INFORMATION:
269 <400> SEQUENCE: 15
                                                                                27
270 atcaaaagat ctgtggctgc accatct
273 <210> SEQ ID NO: 16
274 <211> LENGTH: 58
275 <212> TYPE: DNA
276 <213 > ORGANISM: Artificial Sequence
278 <220> FEATURE:
                             CK1D
279 <223> OTHER INFORMATION:
282 <400> SEQUENCE: 16
283 gegeegteta gaattaacac teteceetgt tgaagetett tgtgaeggge gaacteag
                                                                                58
286 <210> SEQ ID NO: 17
287 <211> LENGTH: 27
288 <212> TYPE: DNA
289 <213> ORGANISM: Artificial Sequence
291 <220> FEATURE:
292 <223> OTHER INFORMATION: (YM001N
295 <400> SEQUENCE: 17
                                                                                27
296 ccggaattca cattcacgat gtacttg
299 <210> SEQ ID NO: 18
300 <211> LENGTH: 16
301 <212> TYPE: DNA
302 <213> ORGANISM: Artificial Sequence
304 <220> FEATURE:
                             YM003
305 <223> OTHER INFORMATION:
308 <400> SEQUENCE: 18
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/508,759

DATE: 09/29/2004 TIME: 16:59:56

Input Set : A:\30215APG\_SEQ.txt

309	tgccccaga ggtgct	16
312	<210> SEQ ID NO: 19	
313	<211> LENGTH: 33	
314	<212> TYPE: DNA	
315	<213> ORGANISM: Artificial Sequence	
317	<220> FEATURE:	
	<223> OTHER INFORMATION (ym257)	
	<400> SEQUENCE: 19	
	acgcattcag tgcttcttgg atgaactggg tga	33
	<210> SEQ ID NO: 20	
	<211> LENGTH: 31	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION (YM258)	
	<400> SEQUENCE: 20	
	atccaagaag cactgaatgc gtagccagaa g	31
	<210> SEQ ID NO: 21	<b>J L</b>
	<211> LENGTH: 38	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence <220> FEATURE:	
	<223> OTHER INFORMATION: YM004	
	<400> SEQUENCE: 21	38
	ccaattcaaa geggttttte cattactata taagagge	30
	<210> SEQ ID NO: 22	
	<211> LENGTH: 32	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION (YM009)	
	<400> SEQUENCE: 22	2.2
	gcagccaccg tacgtttgat ttccaccttg gt	32
	<210> SEQ ID NO: 23	
	<211> LENGTH: 39	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Ryu 166	
	<400> SEQUENCE: 23	
	ggatttgtct gcagtcattg tggctctgcc ctggaactt	39
	<210> SEQ ID NO: 24	
	<211> LENGTH: 27	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Hur 37	
	<400> SEQUENCE: 24	
387	gacaaatcca cgagcacagt ctacatg	27
	JUL TUM!	
	seguer)	
	<220> FEATURE: <223> OTHER INFORMATION: Hur 37 <400> SEQUENCE: 24 gacaaatcca cgagcacagt ctacatg  Llis Hypl  MM in subsequent	

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/508,759

DATE: 09/29/2004

TIME: 16:59:57

Input Set : A:\30215APG\_SEQ.txt

Output Set: N:\CRF4\09292004\J508759.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date